



SEQUENCE LISTING

<110> Larenas, Edmund A.
Goedegebuur, Frits
Gualfetti, Peter
Mitchinson, Colin

<120> Variant Humicola grisea CBH1.1

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<140> US 10/810,277

<141> 2004-03-26

<150> US 60/459,734

<151> 2003-04-01

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<213> Humicola grisea

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<213> Humicola grisea

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<212> PRT

<213> Humicola grisea

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Gly	Ala	Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg
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<212> PRT

<213> Humicola grisea

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Thr	Leu	Asp	Ser	Asn	Trp	Arg	Trp	Thr	His	Gln	Val	Ser	Gly	Ser	Thr		
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<212> PRT

<213> Humicola grisea

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Gly	Asp	Ser	Cys	Gly	Gly	Thr	Tyr	Ser	Asn	Glu	Arg	Tyr	Ala	Gly	Val	
				245				250						255		
Cys	Asp	Pro	Asp	Gly	Cys	Asp	Phe	Asn	Ser	Tyr	Arg	Gln	Gly	Asn	Lys	
			260					265					270			
Thr	Phe	Tyr	Gly	Lys	Gly	Met	Thr	Val	Asp	Thr	Thr	Lys	Lys	Ile	Thr	
		275					280					285				
Val	Val	Thr	Gln	Phe	Leu	Lys	Asp	Ala	Asn	Gly	Asp	Leu	Gly	Glu	Ile	
		290				295					300					
Lys	Arg	Phe	Tyr	Val	Gln	Asp	Gly	Lys	Ile	Ile	Pro	Asn	Ser	Glu	Ser	
305					310					315					320	
Thr	Ile	Pro	Gly	Val	Glu	Gly	Asn	Ser	Ile	Thr	Gln	Asp	Trp	Cys	Asp	
				325				330						335		
Arg	Gln	Lys	Val	Ala	Phe	Gly	Asp	Ile	Asp	Asp	Phe	Asn	Arg	Lys	Gly	
			340					345					350			
Gly	Met	Lys	Gln	Met	Gly	Lys	Ala	Leu	Ala	Gly	Pro	Met	Val	Leu	Val	
		355					360					365				
Met	Ser	Ile	Trp	Asp	Asp	His	Ala	Ser	Asn	Met	Leu	Trp	Leu	Asp	Ser	
		370				375					380					
Thr	Phe	Pro	Val	Asp	Ala	Ala	Gly	Lys	Pro	Gly	Ala	Glu	Arg	Gly	Ala	
385					390					395					400	
Cys	Pro	Thr	Thr	Ser	Gly	Val	Pro	Ala	Glu	Val	Glu	Ala	Glu	Ala	Pro	
				405					410					415		
Asn	Ser	Asn	Val	Val	Phe	Ser	Asn	Ile	Arg	Phe	Gly	Pro	Ile	Gly	Ser	
			420					425					430			
Thr	Val	Ala	Gly	Leu	Pro	Gly	Ala	Gly	Asn	Gly	Gly	Asn	Asn	Gly	Gly	
		435					440					445				
Asn	Pro	Pro	Pro	Thr	Thr	Thr	Thr	Ser	Ser	Ala	Pro	Ala	Thr	Thr		
		450				455				460						
Thr	Thr	Ala	Ser	Ala	Gly	Pro	Lys	Ala	Gly	Arg	Trp	Gln	Gln	Cys	Gly	
465					470					475					480	
Gly	Ile	Gly	Phe	Thr	Gly	Pro	Thr	Gln	Cys	Glu	Glu	Pro	Tyr	Thr	Cys	
				485					490					495		
Thr	Lys	Leu	Asn	Asp	Trp	Tyr	Ser	Gln	Cys	Leu						

<210> 7
 <211> 1662
 <212> DNA
 <213> *Scytalidium thermophilum*

<400> 7
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 gcgtgcagcc tcaccaccga gaggcaccct tccctctcct ggaagaagtg caccgCCggc 120
 ggccagtGCC agaccgtcca ggcttccatc actctcgact ccaactggcg ctggactcac 180
 caggtgtctg gctccaccaa ctgctacacg ggcaacgagt gggattctag catctgcact 240
 gatGCCaagt cgtgcgctca gaactgctgc gtcgatgggtg ctgactacac cagcacctat 300
 ggcatcacca ccaacgggtga ttccctgagc ctcaagttcg tcaccaaggg ccagtactcg 360
 accaacgtcg gctcgcgtac ctacctgatg gacggcgagg acaagtatca gagtaggttc 420
 tatcttcagc cttctcgCGc cttgaatcct ggctaacttt tacacttcac agccttcgag 480
 ctccctcgGca acgagttcac cttcgatgtc gatgtctcca acatcggtcg cgggtctcaac 540
 ggCGccctgt acttcgtctc catggacgcc gatgggtggtc tcagccgcta tcctggcaac 600
 aaggctgggtg ccaagtacgg taccggctac tgcgatgctc agtgcccccg tgacatcaag 660
 ttcatcaacg gcgaggccaa cattgagggc tggaccggct ccaccaacga cccaacgcc 720
 ggCGcgggcc gctatgggtac ctgctgctct gagatggata tctgggaggc caacaacatg 780
 gctactgcct tcactcctca cccttgcaact atcattggcc agagccgctg cgagggcgac 840
 tcgtgcgggtg gcacctacag caacgacCGc tacgCCggcg tctgcgaccc cgatggctgc 900
 gacttcaacg cgtatcgcca gggcaacaag accttctacg gcaagggcat gaccgtcgac 960
 accaccaaga agctcaccgt cgtcaccCGc ttccctcaagg acgccaacgg cgatctcgGC 1020
 gagatcaagc gcttctacgt ccaggatggg aagatcatcc ccaactccga gtccaccatc 1080
 cccggcgctcg agggcaactc catcaccCGc gattgggtgCG accGCCagaa ggttgccctt 1140
 ggCGacattg acgacttcaa ccgcaaggGC ggcatgaagc agatgggcaa ggccctcgcc 1200
 ggccccatgg tccctggctcat gtccatctgg gatgaccag cctccaacat gctctggctc 1260
 gactCGacct tccctgtCGa tgccgctggc aagccCGcg cCGagcgCGg tgctgcccG 1320
 accacctCGg gtgtccctgc tgaggttgag gccgaggccc ccaacagcaa cgtcgtcttc 1380
 tccaacatcc gcttcggccc catcggtcCG accgttgccg gccttcccag cgatggcgGC 1440
 aacaacggcg gcaacaccac cgtccagccc ccgcccagca ccaccaccac ctctgccagc 1500
 agcagcacca cctcggtcc tgccaccacc accaccGCCa gcgctggccc caaggctggc 1560
 cgctggcagc agtgCGggcg catcggtctc actggcccga cccagtgcga ggagccctac 1620
 acttgcacca agctcaacga ctggtactct cagtgcctgt aa 1662

<210> 8
 <211> 1602
 <212> DNA
 <213> *Scytalidium thermophilum*

<400> 8
 atgCGTaccg ccaagttcgc caccctcgcc gcccttgtgg cctcgGCCgc cgcccagcag 60
 gcgtgcagcc tcaccaccga gaggcaccct tccctctcct ggaagaagtg caccgCCggc 120
 ggccagtGCC agaccgtcca ggcttccatc actctcgact ccaactggcg ctggactcac 180
 caggtgtctg gctccaccaa ctgctacacg ggcaacgagt gggattctag catctgcact 240
 gatGCCaagt cgtgcgctca gaactgctgc gtcgatgggtg ctgactacac cagcacctat 300
 ggcatcacca ccaacgggtga ttccctgagc ctcaagttcg tcaccaaggg ccagtactcg 360
 accaacgtcg gctcgcgtac ctacctgatg gacggcgagg acaagtatca gaccttcgag 420
 ctccctcgGca acgagttcac cttcgatgtc gatgtctcca acatcggtcg cgggtctcaac 480
 ggCGccctgt acttcgtctc catggacgcc gatgggtggtc tcagccgcta tcctggcaac 540
 aaggctgggtg ccaagtacgg taccggctac tgcgatgctc agtgcccccg tgacatcaag 600
 ttcatcaacg gcgaggccaa cattgagggc tggaccggct ccaccaacga cccaacgcc 660
 ggCGcgggcc gctatgggtac ctgctgctct gagatggata tctgggaggc caacaacatg 720
 gctactgcct tcactcctca cccttgcaact atcattggcc agagccgctg cgagggcgac 780
 tcgtgcgggtg gcacctacag caacgacCGc tacgCCggcg tctgcgaccc cgatggctgc 840
 gacttcaacg cgtatcgcca gggcaacaag accttctacg gcaagggcat gaccgtcgac 900
 accaccaaga agctcaccgt cgtcaccCGc ttccctcaagg acgccaacgg cgatctcgGC 960
 gagatcaagc gcttctacgt ccaggatggg aagatcatcc ccaactccga gtccaccatc 1020
 cccggcgctcg agggcaactc catcaccCGc gattgggtgCG accGCCagaa ggttgccctt 1080
 ggCGacattg acgacttcaa ccgcaaggGC ggcatgaagc agatgggcaa ggccctcgcc 1140

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ggcccatgg tcctggtcat gtccatctgg gatgaccacg cctccaacat gctctggctc 1200
gactcgacct tccctgtcga tgccgtggc aagcccgcg ccgagcgcg tgccctgccg 1260
accacctcgg gtgtccctgc tgaggttgag gccgaggccc ccaacagcaa cgctcgtctc 1320
tccaacatcc gcttcggccc catcggtcgc accgttgccg gccttcccag cgatggcggc 1380
aacaacggcg gcaacaccac cgtccagccc ccgcccagca ccaccaccac ctctgccagc 1440
agcagcacca cctcgggtcc tgccaccacc accaccgcca gcgctggccc caaggctggc 1500
cgctggcagc agtgcgggcg catcggttc actggcccga cccagtgcga ggagccctac 1560
acttgcacca agctcaacga ctggtactct cagtgcctgt aa 1602

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<210> 9

<211> 533

<212> PRT

<213> Scytalidium thermophilum

<400> 9

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Met Arg Thr Ala Lys Phe Ala Thr Leu Ala Ala Leu Val Ala Ser Ala
1      5      10      15
Ala Ala Gln Gln Ala Cys Ser Leu Thr Thr Glu Arg His Pro Ser Leu
20      25      30
Ser Trp Lys Lys Cys Thr Ala Gly Gly Gln Cys Gln Thr Val Gln Ala
35      40      45
Ser Ile Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly
50      55      60
Ser Thr Asn Cys Tyr Thr Gly Asn Glu Trp Asp Ser Ser Ile Cys Thr
65      70      75      80
Asp Ala Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr
85      90      95
Thr Ser Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys
100     105     110
Phe Val Thr Lys Gly Gln Tyr Ser Thr Asn Val Gly Ser Arg Thr Tyr
115     120     125
Leu Met Asp Gly Glu Asp Lys Tyr Gln Thr Phe Glu Leu Leu Gly Asn
130     135     140
Glu Phe Thr Phe Asp Val Asp Val Ser Asn Ile Gly Cys Gly Leu Asn
145     150     155     160
Gly Ala Leu Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Leu Ser Arg
165     170     175
Tyr Pro Gly Asn Lys Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp
180     185     190
Ala Gln Cys Pro Arg Asp Ile Lys Phe Ile Asn Gly Glu Ala Asn Ile
195     200     205
Glu Gly Trp Thr Gly Ser Thr Asn Asp Pro Asn Ala Gly Ala Gly Arg
210     215     220
Tyr Gly Thr Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Asn Met
225     230     235     240
Ala Thr Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg
245     250     255
Cys Glu Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Asp Arg Tyr Ala
260     265     270
Gly Val Cys Asp Pro Asp Gly Cys Asp Phe Asn Ala Tyr Arg Gln Gly
275     280     285
Asn Lys Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys
290     295     300
Leu Thr Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly
305     310     315     320
Glu Ile Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser
325     330     335
Glu Ser Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp
340     345     350
Cys Asp Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg
355     360     365
Lys Gly Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val

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370	375	380
Leu Val Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu		
385	390	395
Asp Ser Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg		400
	405	410
Gly Ala Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu		415
	420	425
Ala Pro Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile		430
	435	440
Gly Ser Thr Val Ala Gly Leu Pro Ser Asp Gly Gly Asn Asn Gly Gly		445
	450	455
Asn Thr Thr Val Gln Pro Pro Pro Ser Thr Thr Thr Thr Ser Ala Ser		460
465	470	475
Ser Ser Thr Thr Ser Ala Pro Ala Thr Thr Thr Thr Ala Ser Ala Gly		480
	485	490
Pro Lys Ala Gly Arg Trp Gln Gln Cys Gly Gly Ile Gly Phe Thr Gly		495
	500	505
Pro Thr Gln Cys Glu Glu Pro Tyr Thr Cys Thr Lys Leu Asn Asp Trp		510
	515	520
Tyr Ser Gln Cys Leu		525
530		

<210> 10

<211> 497

<212> PRT

<213> Hypocrea jecorina

<400> 10

Gln Ser Ala Cys Thr Leu Gln Ser Glu Thr His Pro Pro Leu Thr Trp		
1	5	10
Gln Lys Cys Ser Ser Gly Gly Thr Cys Thr Gln Gln Thr Gly Ser Val		15
	20	25
Val Ile Asp Ala Asn Trp Arg Trp Thr His Ala Thr Asn Ser Ser Thr		30
	35	40
Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn		45
50	55	60
Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser		65
65	70	75
Thr Tyr Gly Val Thr Thr Ser Gly Asn Ser Leu Ser Ile Gly Phe Val		80
	85	90
Thr Gln Ser Ala Gln Lys Asn Val Gly Ala Arg Leu Tyr Leu Met Ala		95
	100	105
Ser Asp Thr Thr Tyr Gln Glu Phe Thr Leu Leu Gly Asn Glu Phe Ser		110
	115	120
Phe Asp Val Asp Val Ser Gln Leu Pro Cys Gly Leu Asn Gly Ala Leu		125
	130	135
Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Val Ser Lys Tyr Pro Thr		140
145	150	155
Asn Thr Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ser Gln Cys		160
	165	170
Pro Arg Asp Leu Lys Phe Ile Asn Gly Gln Ala Asn Val Glu Gly Trp		175
	180	185
Glu Pro Ser Ser Asn Asn Ala Asn Thr Gly Ile Gly Gly His Gly Ser		190
	195	200
Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Ser Ile Ser Glu Ala		205
	210	215
Leu Thr Pro His Pro Cys Thr Thr Val Gly Gln Glu Ile Cys Glu Gly		220
225	230	235
Asp Gly Cys Gly Gly Thr Tyr Ser Asp Asn Arg Tyr Gly Gly Thr Cys		240
	245	250
Asp Pro Asp Gly Cys Asp Trp Asn Pro Tyr Arg Leu Gly Asn Thr Ser		255
	260	265
		270

Phe	Tyr	Gly	Pro	Gly	Ser	Ser	Phe	Thr	Leu	Asp	Thr	Thr	Lys	Lys	Leu
		275					280					285			
Thr	Val	Val	Thr	Gln	Phe	Glu	Thr	Ser	Gly	Ala	Ile	Asn	Arg	Tyr	Tyr
	290					295					300				
Val	Gln	Asn	Gly	Val	Thr	Phe	Gln	Gln	Pro	Asn	Ala	Glu	Leu	Gly	Ser
305					310					315					320
Tyr	Ser	Gly	Asn	Glu	Leu	Asn	Asp	Asp	Tyr	Cys	Thr	Ala	Glu	Glu	Ala
			325						330					335	
Glu	Phe	Gly	Gly	Ser	Ser	Phe	Ser	Asp	Lys	Gly	Gly	Leu	Thr	Gln	Phe
			340					345					350		
Lys	Lys	Ala	Thr	Ser	Gly	Gly	Met	Val	Leu	Val	Met	Ser	Leu	Trp	Asp
		355					360					365			
Asp	Tyr	Tyr	Ala	Asn	Met	Leu	Trp	Leu	Asp	Ser	Thr	Tyr	Pro	Thr	Asn
	370					375					380				
Glu	Thr	Ser	Ser	Thr	Pro	Gly	Ala	Val	Arg	Gly	Ser	Cys	Ser	Thr	Ser
385					390					395					400
Ser	Gly	Val	Pro	Ala	Gln	Val	Glu	Ser	Gln	Ser	Pro	Asn	Ala	Lys	Val
				405					410					415	
Thr	Phe	Ser	Asn	Ile	Lys	Phe	Gly	Pro	Ile	Gly	Ser	Thr	Gly	Asn	Pro
			420					425					430		
Ser	Gly	Gly	Asn	Pro	Pro	Gly	Gly	Asn	Pro	Pro	Gly	Thr	Thr	Thr	Thr
			435				440					445			
Arg	Arg	Pro	Ala	Thr	Thr	Thr	Gly	Ser	Ser	Pro	Gly	Pro	Thr	Gln	Ser
	450					455					460				
His	Tyr	Gly	Gln	Cys	Gly	Gly	Ile	Gly	Tyr	Ser	Gly	Pro	Thr	Val	Cys
465					470					475					480
Ala	Ser	Gly	Thr	Thr	Cys	Gln	Val	Leu	Asn	Pro	Tyr	Tyr	Ser	Gln	Cys
				485					490					495	
Leu															

<210> 11

<211> 515

<212> PRT

<213> Scytalidium thermophilum

<400> 11

Gln	Gln	Ala	Cys	Ser	Leu	Thr	Thr	Glu	Arg	His	Pro	Ser	Leu	Ser	Trp
1				5					10					15	
Lys	Lys	Cys	Thr	Ala	Gly	Gly	Gln	Cys	Gln	Thr	Val	Gln	Ala	Ser	Ile
			20					25					30		
Thr	Leu	Asp	Ser	Asn	Trp	Arg	Trp	Thr	His	Gln	Val	Ser	Gly	Ser	Thr
	35						40					45			
Asn	Cys	Tyr	Thr	Gly	Asn	Glu	Trp	Asp	Ser	Ser	Ile	Cys	Thr	Asp	Ala
50						55					60				
Lys	Ser	Cys	Ala	Gln	Asn	Cys	Cys	Val	Asp	Gly	Ala	Asp	Tyr	Thr	Ser
65					70					75					80
Thr	Tyr	Gly	Ile	Thr	Thr	Asn	Gly	Asp	Ser	Leu	Ser	Leu	Lys	Phe	Val
				85					90					95	
Thr	Lys	Gly	Gln	Tyr	Ser	Thr	Asn	Val	Gly	Ser	Arg	Thr	Tyr	Leu	Met
			100					105					110		
Asp	Gly	Glu	Asp	Lys	Tyr	Gln	Thr	Phe	Glu	Leu	Leu	Gly	Asn	Glu	Phe
		115					120					125			
Thr	Phe	Asp	Val	Asp	Val	Ser	Asn	Ile	Gly	Cys	Gly	Leu	Asn	Gly	Ala
	130					135					140				
Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg	Tyr	Pro
145					150					155					160
Gly	Asn	Lys	Ala	Gly	Ala	Lys	Tyr	Gly	Thr	Gly	Tyr	Cys	Asp	Ala	Gln
				165					170					175	
Cys	Pro	Arg	Asp	Ile	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Ile	Glu	Gly
			180					185					190		
Trp	Thr	Gly	Ser	Thr	Asn	Asp	Pro	Asn	Ala	Gly	Ala	Gly	Arg	Tyr	Gly

195	200	205
Thr Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Asn Met Ala Thr		
210	215	220
Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg Cys Glu		
225	230	235
Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Asp Arg Tyr Ala Gly Val		
245	250	255
Cys Asp Pro Asp Gly Cys Asp Phe Asn Ala Tyr Arg Gln Gly Asn Lys		
260	265	270
Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys Leu Thr		
275	280	285
Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly Glu Ile		
290	295	300
Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser Glu Ser		
305	310	315
Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp Cys Asp		
325	330	335
Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg Lys Gly		
340	345	350
Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val Leu Val		
355	360	365
Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu Asp Ser		
370	375	380
Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg Gly Ala		
385	390	395
Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu Ala Pro		
405	410	415
Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile Gly Ser		
420	425	430
Thr Val Ala Gly Leu Pro Ser Asp Gly Gly Asn Asn Gly Gly Asn Thr		
435	440	445
Thr Val Gln Pro Pro Pro Ser Thr Thr Thr Thr Ser Ala Ser Ser Ser		
450	455	460
Thr Thr Ser Ala Pro Ala Thr Thr Thr Thr Ala Ser Ala Gly Pro Lys		
465	470	475
Ala Gly Arg Trp Gln Gln Cys Gly Gly Ile Gly Phe Thr Gly Pro Thr		
485	490	495
Gln Cys Glu Glu Pro Tyr Thr Cys Thr Lys Leu Asn Asp Trp Tyr Ser		
500	505	510
Gln Cys Leu		
515		

<210> 12
 <211> 507
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> consensus sequence

<400> 12

Gln Gln Ala Cys Ser Leu Thr Thr Glu Arg His Pro Ser Leu Ser Trp
1 5 10 15
Lys Lys Cys Thr Ala Gly Gly Gln Cys Gln Thr Val Gln Ala Ser Ile
20 25 30
Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly Ser Thr
35 40 45
Asn Cys Tyr Thr Gly Asn Lys Trp Asp Ser Ser Ile Cys Thr Asp Ala
50 55 60
Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr Thr Ser
65 70 75 80
Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys Phe Val

				85					90					95			
Thr	Lys	Gly	Gln	His	Ser	Thr	Asn	Val	Gly	Ser	Arg	Thr	Tyr	Leu	Met		
			100					105					110				
Asp	Gly	Glu	Asp	Lys	Tyr	Gln	Thr	Phe	Glu	Leu	Leu	Gly	Asn	Glu	Phe		
		115				120						125					
Thr	Phe	Asp	Val	Asp	Val	Ser	Asn	Ile	Gly	Cys	Gly	Leu	Asn	Gly	Ala		
	130					135					140						
Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg	Tyr	Pro		
145					150					155					160		
Gly	Asn	Lys	Ala	Gly	Ala	Lys	Tyr	Gly	Thr	Gly	Tyr	Cys	Asp	Ala	Gln		
			165					170						175			
Cys	Pro	Arg	Asp	Ile	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Ile	Glu	Gly		
		180						185					190				
Trp	Thr	Gly	Ser	Thr	Asn	Asp	Pro	Asn	Ala	Gly	Ala	Gly	Arg	Tyr	Gly		
	195					200						205					
Thr	Cys	Cys	Ser	Glu	Met	Asp	Ile	Trp	Glu	Ala	Asn	Asn	Met	Ala	Thr		
	210					215						220					
Ala	Phe	Thr	Pro	His	Pro	Cys	Thr	Ile	Ile	Gly	Gln	Ser	Arg	Cys	Glu		
225					230					235					240		
Gly	Asp	Ser	Cys	Gly	Gly	Thr	Tyr	Ser	Asn	Glu	Arg	Tyr	Ala	Gly	Val		
			245					250						255			
Cys	Asp	Pro	Asp	Gly	Cys	Asp	Phe	Asn	Ser	Tyr	Arg	Gln	Gly	Asn	Lys		
		260						265					270				
Thr	Phe	Tyr	Gly	Lys	Gly	Met	Thr	Val	Asp	Thr	Thr	Lys	Lys	Ile	Thr		
	275						280					285					
Val	Val	Thr	Gln	Phe	Leu	Lys	Asp	Ala	Asn	Gly	Asp	Leu	Gly	Glu	Ile		
	290					295					300						
Lys	Arg	Phe	Tyr	Val	Gln	Asp	Gly	Lys	Ile	Ile	Pro	Asn	Ser	Glu	Ser		
305					310					315					320		
Thr	Ile	Pro	Gly	Val	Glu	Gly	Asn	Ser	Ile	Thr	Gln	Asp	Trp	Cys	Asp		
			325					330						335			
Arg	Gln	Lys	Val	Ala	Phe	Gly	Asp	Ile	Asp	Asp	Phe	Asn	Arg	Lys	Gly		
		340						345					350				
Gly	Met	Lys	Gln	Met	Gly	Lys	Ala	Leu	Ala	Gly	Pro	Met	Val	Leu	Val		
	355						360					365					
Met	Ser	Ile	Trp	Asp	Asp	His	Ala	Ser	Asn	Met	Leu	Trp	Leu	Asp	Ser		
	370					375					380						
Thr	Phe	Pro	Val	Asp	Ala	Ala	Gly	Lys	Pro	Gly	Ala	Glu	Arg	Gly	Ala		
385					390					395					400		
Cys	Pro	Thr	Thr	Ser	Gly	Val	Pro	Ala	Glu	Val	Glu	Ala	Glu	Ala	Pro		
			405					410						415			
Asn	Ser	Asn	Val	Val	Phe	Ser	Asn	Ile	Arg	Phe	Gly	Pro	Ile	Gly	Ser		
		420						425					430				
Thr	Val	Ala	Gly	Leu	Pro	Gly	Ala	Gly	Asn	Gly	Gly	Asn	Asn	Gly	Gly		
	435					440						445					
Asn	Pro	Pro	Pro	Pro	Thr	Thr	Thr	Thr	Ser	Ser	Ala	Pro	Ala	Thr	Thr		
	450					455					460						
Thr	Thr	Ala	Ser	Ala	Gly	Pro	Lys	Ala	Gly	Arg	Trp	Gln	Gln	Cys	Gly		
465					470					475					480		
Gly	Ile	Gly	Phe	Thr	Gly	Pro	Thr	Gln	Cys	Glu	Glu	Pro	Tyr	Thr	Cys		
			485					490						495			
Thr	Lys	Leu	Asn	Asp	Trp	Tyr	Ser	Gln	Cys	Leu							
		500						505									

<210> 13

<211> 507

<212> PRT

<213> Artificial Sequence

<220>

<223> consensus sequence

<400> 13

Gln	Gln	Ala	Cys	Ser	Leu	Thr	Thr	Glu	Arg	His	Pro	Ser	Leu	Ser	Trp
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Lys	Lys	Cys	Thr	Ala	Gly	Gly	Gln	Cys	Gln	Thr	Val	Gln	Ala	Ser	Ile
		20					25					30			
Thr	Leu	Asp	Ser	Asn	Trp	Arg	Trp	Thr	His	Gln	Val	Ser	Gly	Ser	Thr
	35					40					45				
Asn	Cys	Tyr	Thr	Gly	Asn	Lys	Trp	Asp	Thr	Ser	Ile	Cys	Thr	Asp	Ala
	50				55						60				
Lys	Ser	Cys	Ala	Gln	Asn	Cys	Cys	Val	Asp	Gly	Ala	Asp	Tyr	Thr	Ser
65				70					75					80	
Thr	Tyr	Gly	Ile	Thr	Thr	Asn	Gly	Asp	Ser	Leu	Ser	Leu	Lys	Phe	Val
			85					90						95	
Thr	Lys	Gly	Gln	His	Ser	Thr	Asn	Val	Gly	Ser	Arg	Thr	Tyr	Leu	Met
		100					105						110		
Asp	Gly	Glu	Asp	Lys	Tyr	Gln	Thr	Phe	Glu	Leu	Leu	Gly	Asn	Glu	Phe
	115						120					125			
Thr	Phe	Asp	Val	Asp	Val	Ser	Asn	Ile	Gly	Cys	Gly	Leu	Asn	Gly	Ala
	130					135					140				
Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg	Tyr	Pro
145				150					155						160
Gly	Asn	Lys	Ala	Gly	Ala	Lys	Tyr	Gly	Thr	Gly	Tyr	Cys	Asp	Ala	Gln
			165					170						175	
Cys	Pro	Arg	Asp	Ile	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Ile	Glu	Gly
		180					185					190			
Trp	Thr	Gly	Ser	Thr	Asn	Asp	Pro	Asn	Ala	Gly	Ala	Gly	Arg	Tyr	Gly
	195					200						205			
Thr	Cys	Cys	Ser	Glu	Met	Asp	Ile	Trp	Glu	Ala	Asn	Asn	Met	Ala	Thr
	210				215						220				
Ala	Phe	Thr	Pro	His	Pro	Cys	Thr	Ile	Ile	Gly	Gln	Ser	Arg	Cys	Glu
225				230					235						240
Gly	Asp	Ser	Cys	Gly	Gly	Thr	Tyr	Ser	Asn	Glu	Arg	Tyr	Ala	Gly	Val
			245					250						255	
Cys	Asp	Pro	Asp	Gly	Cys	Asp	Phe	Asn	Ser	Tyr	Arg	Gln	Gly	Asn	Lys
		260					265					270			
Thr	Phe	Tyr	Gly	Lys	Gly	Met	Thr	Val	Asp	Thr	Thr	Lys	Lys	Ile	Thr
	275					280						285			
Val	Val	Thr	Gln	Phe	Leu	Lys	Asp	Ala	Asn	Gly	Asp	Leu	Gly	Glu	Ile
	290				295						300				
Lys	Arg	Phe	Tyr	Val	Gln	Asp	Gly	Lys	Ile	Ile	Pro	Asn	Ser	Glu	Ser
305				310					315						320
Thr	Ile	Pro	Gly	Val	Glu	Gly	Asn	Ser	Ile	Thr	Gln	Asp	Trp	Cys	Asp
			325					330						335	
Arg	Gln	Lys	Val	Ala	Phe	Gly	Asp	Ile	Asp	Asp	Phe	Asn	Arg	Lys	Gly
		340					345					350			
Gly	Met	Lys	Gln	Met	Gly	Lys	Ala	Leu	Ala	Gly	Pro	Met	Val	Leu	Val
	355				360						365				
Met	Ser	Ile	Trp	Asp	Asp	His	Ala	Ser	Asn	Met	Leu	Trp	Leu	Asp	Ser
	370				375					380					
Thr	Phe	Pro	Val	Asp	Ala	Ala	Gly	Lys	Pro	Gly	Ala	Glu	Arg	Gly	Ala
385				390					395						400
Cys	Pro	Thr	Thr	Ser	Gly	Val	Pro	Ala	Glu	Val	Glu	Ala	Glu	Ala	Pro
			405					410					415		
Asn	Ser	Asn	Val	Phe	Ser	Asn	Ile	Arg	Phe	Gly	Pro	Ile	Gly	Ser	
		420					425					430			
Thr	Val	Ala	Gly	Leu	Pro	Gly	Ala	Gly	Asn	Gly	Gly	Asn	Asn	Gly	Gly
	435					440					445				
Asn	Pro	Pro	Pro	Pro	Thr	Thr	Thr	Thr	Ser	Ser	Ala	Pro	Ala	Thr	Thr
	450				455						460				
Thr	Thr	Ala	Ser	Ala	Gly	Pro	Lys	Ala	Gly	Arg	Trp	Gln	Gln	Cys	Gly
465				470					475						480
Gly	Ile	Gly	Phe	Thr	Gly	Pro	Thr	Gln	Cys	Glu	Glu	Pro	Tyr	Thr	Cys

